## CLAIMS :

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- 1. A cloned DNA which contains a DNA which is hybridizable with the genomic RNA of the LAV virues or a fragment of said hybridizable DNA.
- 2. The DNA for claim t which is a recombinent of said hybridizable DNA or DNA fragment hybridizable with the conduct RNA of the LAV virus.
- 3. The DNA of claim 1 or 2 wherein said hybradicable DNA or DNA fragment is a cDNA.
- '4. The DNA of claims 1 to 3 which contains the following restriction sites in the following order (from the 3' and to the 5' end):

Hind III, Sac I, Cgl II (LAV 75).

5. The DHA of claim 4 which contains the following restriction sites in the following order:

Hind III, Sac I, Byl II, Byl II, Kpm I (LAV 82).

6. The DNA of claim witch contains the following restriction sites in the following order:

Hind III, Sec I, Fol II, Bol II, Kon I, XHO I, Bom HI, Hind III, Bol II (LAV 13).

7. The DNA of claim 6 which has a size of about 5 kb.

- C. The DHA of any of claims 1 to 7 which contains a region corresponding to the R and U3 regions of the LTR as well as to the 3' end of the coding region of the retroviral DNA.
- 9. The DNA of claim I which has a size from about 9.1 to 9.2 kb.
- 10. The DNA of claim 9 which contains the following series of restriction sites:

30	Hind III	0
	Sac I	50
	Bam HI	460
	Hind /II	520
	Bem HI	600
	Pat I	800

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Hind III 1 100 Dgl II 1 50ø Kpn I 5,00 Kon I /9 O O Eco RI 100 Eco RI 5 300 5 5 500 Sal I Kpn I 6 100 8g1 II 6 500 Bgl [II 7 G00 Kind III 7 850 10 - Barn HI 8 150 Xho I 8 600 Kpn I 8 700 Bgl I 8 750 8g1 I 9 150 15 Sac ! 9 200 9 250 Hind VII

11. The DNA of claim 10 which contains or additional Hind III approximately at the 5 550 coordinate.

12. A DNA fragmont according to claim ( which comprises a sequence extending from approximately Kna I (6100) to approximately Bam HI (8150) of the sequence defined in claim 11.

13. A DNA fragment according to claim 1 which comprises a sequence extending from approximately Kon 1 (3500) to approximately Bgl II (6500) of the sequence defined in claim 11.

14. A DNA fragment according to claim 1 which corprises a sequence extending from approximately Pst (800) to approximately Kpn I (3500) of the sequence defined in claim 11.

15. A DNA fragment of claim 1 which codes for the envelopes proteins.

16. A DNA fragment of claim 1 which codes for the retroviral polymerase.

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17. A DHA fragmant which codes proteins.

18. probe for the in vitro detection of LAV which consists of a DNA according to any of claims 1 to

An expression vector, particularly a plasmid, 5 for the transformation of procaryotic or sucaryotic cells which contains an insert consisting of a DNA fragmont hybridizable with the retroviral genome of LAV viruses as defined in any of claims 1 to 17.

20. The vector of claim 18 which contains the DNA fragment of claim 15.

21. A microorganism, eucaryotic or procaryotic coll which is transformed by a vector according to claim 19 or 20 and which expresses the polypeptide encoded by the corresponding DNA fragment,

22. The purified RNAs of LAV viruses which have sizes from 9.1 to 9.2 kb.

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